



STIC Search Report

Biotech-Chem Library

STIC Database Tracking Number: 179017

TO: Christian Fronda
Location: REM-2D78/2C70
Art Unit: 1652
Wednesday, February 22, 2006
Case Serial Number: 10/043639

From: Barb O'Bryen
Location: Biotech-Chem Library
Remsen 1a69
Phone: 571-272-2518
POB
barbara.obryen@uspto.gov

Search Notes

O'Bryen, Barbara

From: Fronda, Christian
Sent: Wednesday, February 15, 2006 4:43 PM
To: O'Bryen, Barbara
Subject: RE: problem with Sequence search for Serial No. 10/043,639

Importance: High

Can you search only SEQ ID NOS: 3, 5, 6, 7, and 8. Please advise if the search exceeds the 20hrs limit. Thank you very much.

Christian L. Fronda
Art Unit 1652
Office REM 2D78
Mailbox REM 2C70
(571)272-0929

8 5 no

lg / default

pwB 2-22-06

①

-----Original Message-----

From: O'Bryen, Barbara
Sent: Wednesday, February 15, 2006 4:27 PM
To: Fronda, Christian
Subject: problem with Sequence search for Serial No. 10/043,639
Importance: High

Ex Fronda,
I'm now working the search you submitted below, but unfortunately the search as requested would take too long to process.

In order to make the most fair use of system resources, the following criteria have been developed for rejecting search requests:

- 1) no more than 10 nucleic acid sequences
- 2) no sequences >9,999 nt long
- 3) total search time must be <20 hrs

The search as requested would require ~28hrs of processing time on our fastest machine.

Please contact one of the examiners listed below, who may be able to help you reduce the size of your request, or can give special authorization to have the search run as is.

Scott Priebe
Jeff Fredman
Jim Martinell
Mike Pak

Please forward the changes or approval directly to me since I have the request on my desk, rather than to the STIC-BioTech/ChemLib mailbox.

Thanks,
Barb O'Bryen
STIC
2-2518

~~SHOTS~~

WTF 8/12/06

-----Original Message-----

From: STIC-BioTech/ChemLib
Sent: Wednesday, February 15, 2006 4:20 PM
To: O'Bryen, Barbara
Subject: FW: Sequence search for Serial No. 10/043,639
Importance: High

-----Original Message-----

From: Fronda, Christian
Sent: Wednesday, February 08, 2006 1:25 PM
To: STIC-Biotech/ChemLib
Subject: Sequence search for Serial No. 10/043,639
Importance: High

Please perform sequence search for Serial No. 10/043,639

Please search SEQ ID Nos.: 1-8 against nucleic acid commercial, PGPub, issued databases, and interference databases.

Please save on COMPUTER DISKETTES.

Please save results from interference data base search on separate and different diskettes.

Thank you very much.

Christian Fronda
Art Unit 1652
Mailbox REM 2C70
Office REM 2D78
(517)272-0929

November 2005

Published_Applications Nucleic Acid and Published_Applications Amino Acid database searches now generate two sets of results each. The Published_Applications databases have been split into two parts to reduce the amount of time required for their daily updates. This results in more machine time being available for processing searches.

Newly published applications will appear in the Published_Applications_New databases; older published applications make up the Published_Applications_Main databases.

Searches run against Nucleic Acid Published_Applications produce two sets of results, with the extensions .rnpbm (Published_Applications_NA_Main) and .rnpbn (Published_Applications_NA_New). Searches run against Amino Acid Published_Applications produce two sets of results, with the extensions .rapbm (Published_Applications_AA_Main) and .rapbn (Published_Applications_AA_New).

Pending Nucleic Acid and Pending Amino Acid database searches generate two sets of results each. The Pending databases have been split into two parts to reduce the amount of time required for their daily updates. This results in more machine time being available for processing searches.

Searches run against the Nucleic Acid Pending database produce two sets of results, with the extensions .rnpm and .rnpn

Searches run against the Amino Acid Pending database produce two sets of results, with the extensions .rapm and .rapn

Because they contain data that is confidential, the results of Pending database searches should not be left in the case.